- 1 ATGCAGAAGA CICTICAAGA TICAGCITIC CIGGAAACIG ATCITCAATG
- 51 CACTAAGAGA AGGAGACTOT CAAACOCAAA AATGACCTGG AGGCACCATG
 M T W R H H
- 101 TCAGGCTTCT GITTACGGTC AGTTTGGCAT TACAGATCAT CAATTTGGCA
 V R L L F T V S L A L O I I N L G
- AACAGCTATC AAAGAGAGAA ACATAACGGC GGTAGAGAGG AAGTCACCAA
- N S Y Q R E K H N G G R E E V T K
 201 GGTTGCCACT CAGAAGCACC GACAGTCACC GCTCAACTGG ACCTCCAGTC
 - VAT QKHR QSP LNW TSS
- 251 ATTTOGGAGA GGTGACTOGG AGGGCGGGGG GCTGGGGGGCCG
- H F G E V T G S A E G W G P E E P
- L P Y S R A F G E G A S A R P R C
- CTGCAGGAAC GGCGGTACCT GCGTGCTGGG CAGCTTCTGC GTGTGCCCCGG
 C R N G G T C V L G S F C V C P
- 401 CCCACTICAC CCCCCCTAC TCCCACCATG ACCACACCC CAGTGAATCC
- A H F T G R Y C E H D Q R R S E C
- 451 GGCGCCCTGG AGCACGGAGC CTGGACCCTC CGCGCCTGCC ACCTCTGCAG
- GALE HGA WTL RACH LCR
- 501 GIGCATCTIC GGGGCCCIGC ACIGCCTCCC CCICCAGACG CCIGACCGCT
 - C I F G A L H C L P L Q T P D R
- 551 GTGACCOGAA AGACTTOCTG GOCTCOCACG CTCACGGGCC GAGCGCCGGG
 - C D P K D F L A S H A H G P S A G
- 601 GGCGCCCCA GCCTGCTACT CTTGCTGCCC TGCGCAACTC CTGCACCGGC
- GAPS LLL LLP CATP APA
- S C A R M R P R T L G P W S L P
- 701 CCTCCAGCGG GAGCGGCGCC CCTGCGGCAAG GCCGGCACTT GGGCATCGCC
- SSSG SGA PAEG RDL GIA
- 751 TITAATITIC TAIGITGIAA ATAA

F N F L C C K

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ICUKE 1

criptin cripto	1 MTWRHHVRLL MDCRKMARFS	FTVSLALQI. YSVIWIMAIS	INLGNSYQ KVFELGLVAG	REKHNGGREE LGHQEFARPS	50 VTKVATQKHR RGYLAFRDDS
criptin cripto	51 QSPLNWTSSH IWPQE	FGEVTGSAEG EPAIRPRSSQ	WGPEEPLPYS RVPPMGIQHS	RAFGEGASAR KELNRT	100 PRCCRNGGTC CCLNGGTC
criptin cripto	101 VLGSFCVCPA MLGSFCACPP	HFTGRYCEHD SFYGRNCEHD	QRRSECGALE VRKENCGSVF	HGAWTLRACH HDTWLPKKCS	150 LCRCIFGALH LCKCWHGQLR
criptin cripto	151 CLPLQTPDRC CFPQAFLPGC	DPKDFL DGLVMDEHLV	ASHAHGPSAG ASRTPELPPS	GAPSLLLLLP ARTTTFMLVG	200 CATPAPASCA ICLSIQSYY.
criptin cripto	201 RMRPRTLGPW	SLPSSSGSGA	PAEGRDLGIA	237 FNFLCCK	

Note: Conserved Cysteine residues are shown in BOLD.

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